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## Cardiac Troponin T Measured by a Highly Sensitive Assay Predicts Coronary Heart Disease, Heart Failure, and Mortality in the ARIC Study

Justin T. Saunders, MD<sup>\*,†</sup>, Vijay Nambi, MD<sup>\*</sup>, James A. de Lemos, MD, Lloyd E. Chambless, PhD, Salim S. Virani, MD, Eric Boerwinkle, PhD, Ron C. Hoogeveen, PhD, Xiaoxi Liu, MS, Brad C. Astor, PhD, Thomas H. Mosley, PhD, Aaron R. Folsom, MD, MPH, Gerardo Heiss, MD, MSc, PhD, Josef Coresh, MD, PhD, and Christie M. Ballantyne, MD  
Department of Medicine, Baylor College of Medicine, and Center for Cardiovascular Disease Prevention, Methodist DeBakey Heart and Vascular Center, Houston, TX (J.T.S., V.N., S.S.V., R.C.H., C.M.B.); Department of Internal Medicine, University of Texas Southwestern Medical Center, Dallas, TX (J.A.D.); Department of Biostatistics, University of North Carolina, Chapel Hill, NC (L.E.C., X.L., G.H.); Michael E. DeBakey Veterans Affairs Medical Center, Houston, TX (S.S.V); Human Genetics Center, University of Texas Health Science Center School of Public Health, Houston, TX (E.B.); Department of Epidemiology, Johns Hopkins Bloomberg School of Public Health, Baltimore, MD (B.C.A., J.C.); Department of Internal Medicine, University of Mississippi Medical Center, Jackson, MS (T.H.M.); and Division of Epidemiology and Community Health, University of Minnesota School of Public Health, Minneapolis, MN (A.R.F)

### Abstract

**Background**—We evaluated whether cardiac troponin T (cTnT) measured with a new highly sensitive assay was associated with incident coronary heart disease (CHD), mortality, and hospitalization for heart failure (HF) in a general population of participants in the Atherosclerosis Risk in Communities (ARIC) Study.

**Methods and Results**—Associations between increasing cTnT levels and CHD, mortality, and HF hospitalization were evaluated using Cox proportional-hazards models adjusted for traditional CHD risk factors, kidney function, high-sensitivity C-reactive protein (hs-CRP), and N-terminal pro-B-type natriuretic peptide (NT-proBNP) in 9,698 participants aged 54–74 years who at baseline were free from CHD and stroke (and HF in the HF analysis). Measurable cTnT levels ( $\geq 0.003$   $\mu\text{g/L}$ ) were detected in 66.5% of individuals. In fully adjusted models, compared with participants with undetectable levels, those with cTnT levels in the highest category ( $\geq 0.014$   $\mu\text{g/L}$ , 7.4% of the ARIC population) had significantly increased risk for CHD (hazard ratio [HR] 2.29, 95% confidence interval [CI] 1.81–2.89), fatal CHD (HR 7.59, 95% CI 3.78–15.25), total mortality (HR 3.96, 95% CI 3.21–4.88), and HF (HR 5.95, 95% CI 4.47–7.92). Even minimally elevated cTnT ( $\geq 0.003$   $\mu\text{g/L}$ ) was associated with increased risk for mortality and HF ( $p < 0.05$ ). Adding cTnT to traditional risk factors improved risk prediction parameters; the improvements were similar to those with NT-proBNP and better than that with the addition of hs-CRP.

Correspondence: Christie M. Ballantyne, MD, 6565 Fannin, M.S. A-601, Houston, TX 77030; phone: 713-798-5034; fax: 713-798-3057; cmb@bcm.tmc.edu.

<sup>\*</sup>Co-primary authors;

<sup>†</sup>Current affiliation: Department of Internal Medicine, University of Michigan, Ann Arbor, MI

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**Conclusions**—cTnT detectable with a highly sensitive assay was associated with incident CHD, mortality, and HF in individuals from a general population without known CHD/stroke.

## Keywords

coronary disease; heart failure; risk factors; troponin

Cardiac troponin T (cTnT) and troponin I are the biomarkers recommended for diagnosis of a myocardial infarction (MI) and in the setting of an acute coronary syndrome.<sup>1</sup> Cardiac troponin, which is released in response to cardiomyocyte necrosis, has been independently associated with adverse outcomes following acute coronary syndrome,<sup>2</sup> in patients with chronic heart failure (HF),<sup>3</sup> and in the general population although cTnT levels are detectable in only a small fraction of the general population using current assays.<sup>4–8</sup>

A precommercial highly sensitive cTnT assay (hs-cTnT) can detect 10-fold lower concentrations than currently available 4th-generation assays. Using this assay, cTnT levels below the detection range of standard assays were independently associated with adverse cardiovascular events in patients with HF<sup>9</sup> or stable coronary heart disease (CHD).<sup>10</sup> The clinical significance of detectable cTnT using the new assay in the general population is not known.

We determined prevalence of measurable cTnT with the new assay; evaluated associations of cTnT with CHD, death, and HF; and determined whether cTnT improves risk prediction for these outcomes in participants without cardiovascular disease (CVD) (n=9,698) in the Atherosclerosis Risk in Communities (ARIC) Study. Further, we performed limited comparisons between cTnT, N-terminal pro-B-type natriuretic peptide (NT-proBNP), and high-sensitivity C-reactive protein (hs-CRP) with respect to risk prediction.

## METHODS

More detailed Methods are described in the Supplementary Material. cTnT levels were measured with a novel precommercial highly sensitive assay (Elecsys Troponin T [Roche Diagnostics, Indianapolis, IN]). The lower limit of detection of the novel assay is 0.003 µg/L. The lower limit of detection of the currently available clinically used assay is 0.01 µg/L, which corresponds to 0.03 µg/L with the high-sensitivity assay (i.e., the new assay has 10-fold increased detection).<sup>10</sup> Outcomes assessed were incident CHD events (fatal CHD, definite or probable MI, coronary revascularization), all-cause mortality, and hospitalization for HF (ICD-9 code 428).

## Statistical Methods

The distribution and percentile of cTnT were determined from all ARIC participants with measurements, including those with prevalent CVD, to reflect more accurately the general population. All other analyses were performed only in participants without CVD (HF analyses also excluded those with prevalent HF; also see Supplementary Material).

We modeled cTnT as both a categorical and a continuous variable. For the categorical analyses, the 33.5% with undetectable levels were the reference group (Group 1). The remaining 66.5% were split into approximate thirds: cTnT levels 0.003–0.005 µg/L (Group 2), 0.006–0.008 µg/L (Group 3), and higher levels divided at approximately the 90<sup>th</sup> percentile of the ARIC population (Group 4: 0.009–0.013 µg/L; Group 5: ≥0.014 µg/L), which incidentally corresponded to the 99<sup>th</sup> percentile value specified by the manufacturer. Because of confounding by age, gender, and race, we present age-, gender-, and race-adjusted survival curves by cTnT level; adjustments were stratified by the 5 cTnT categories

to fit product-limit survival functions for each age/gender/race combination and applying the method of predicted marginals<sup>11</sup> to obtain adjusted survival curves. Age-, gender-, and race-adjusted events rates were estimated by Poisson regression. Associations between cTnT categories and outcomes were determined with Cox proportional-hazards models. The basic model (model 1) adjusted for age, gender, and race. Model 2 adjusted for components of the ARIC CHD Risk Score (ACRS): all components of model 1, total cholesterol, high-density lipoprotein cholesterol (HDL-C), systolic blood pressure, antihypertensive medication use, smoking status, and presence of diabetes (fasting blood glucose >126 mg/dL or antidiabetic medication use).<sup>12</sup> Coefficients for ACRS components were derived specifically for this dataset. Model 3 adjusted for ACRS plus hs-CRP, NT-proBNP, and estimated glomerular filtration rate (eGFR). We also analyzed associations between cTnT and outcomes using a continuous piecewise linear model, which demonstrated a better fit for incident CHD than polynomial or log-transformed models. In continuous models, undetectable levels were assigned a level of 0.0015 µg/L (half the lower limit of detection, as suggested by D'Angelo et al<sup>13</sup>).

To analyze the incremental value of cTnT in risk prediction, areas under the receiver operating characteristic curve (AUC), net reclassification improvement (NRI), clinical NRI (NRI in the intermediate-risk group), and integrated discrimination improvement (IDI) were calculated for the overall study population and by gender for 10-year follow-up. Bootstrapping was performed to furnish 95% confidence intervals (CIs) for the differences between models. Model calibration was assessed by Grønnesby–Borgan goodness-of-fit test. The base model for CHD risk prediction was ACRS alone; the extended model added cTnT as a continuous variable using the piecewise linear model described above. Although no risk prediction models for total mortality and HF hospitalizations are validated in ARIC, we added body mass index (BMI), left ventricular hypertrophy (LVH), and creatinine to ACRS as the base model for prediction of mortality and HF hospitalization as these additional variables were associated with mortality and HF in other studies.<sup>14,15</sup> We created similar risk categories (as for CHD risk prediction) to estimate NRI and clinical NRI. Finally, we added hs-CRP and NT-proBNP to the risk prediction models for CHD, mortality, and HF to compare cTnT with these other markers.

## RESULTS

cTnT was detectable in 6,440 individuals (66.5%) and  $\geq 0.014$  µg/L in 715 individuals (7.4%). The 99<sup>th</sup> percentile for cTnT in the entire ARIC population was 0.03 µg/L (corresponding to the lower limit of detection of the current, clinically used assay) and 0.027 µg/L in participants without CVD. Thus, <1% of the ARIC population had cTnT levels that would have been detected by the currently used assay. Detectable cTnT levels ranged from 0.003 µg/L to 0.340 µg/L. The distribution of cTnT by gender is shown in Figure 1. cTnT was detectable in 83% of men, compared with ~55% of women; 13% of men and 3% of women were in the highest cTnT group. Log-transformed cTnT levels correlated with log-transformed NT-proBNP levels; after excluding individuals with undetectable cTnT and NT-proBNP levels, Pearson's correlation coefficients were 0.22, 0.32, and 0.25 for the overall group, men, and women, respectively ( $p < 0.0001$  for all).

Table 1 shows baseline demographic characteristics by cTnT category. Individuals with higher cTnT levels were older; were more likely to be male, hypertensive, and diabetic; were less likely to be current smokers; had lower eGFR, total cholesterol, and HDL-C; and had higher triglyceride, hs-CRP, and NT-proBNP levels.

## Associations of Troponin Levels with CHD, Mortality, and HF Hospitalization

All hazard ratios (HRs) presented for associations between cTnT and outcomes used the undetectable cTnT group as the referent. Relationships between cTnT and each outcome were not statistically different across time periods (<1 year, 1–3 years, >3 years); therefore, the proportional-hazards assumption was not rejected. Age-, gender-, and race-adjusted event rates are presented in Supplemental Table 1. Overall, event rates increased with increasing cTnT levels, except CHD in men, for which the increase was apparent only in the highest cTnT category.

**CHD**—cTnT values in the highest category were associated with incident CHD events (n=983 [117 CHD deaths, 454 definite or probable MIs, and 916 definite or probable MIs or coronary revascularizations]; mean follow-up 9.4 years) in the overall cohort (Table 2), as well as in men and women (Supplemental Tables 2 and 3) considered separately. Associations of cTnT with fatal CHD (HR 7.59, 95% CI 3.78–15.25 for Group 5) were consistently stronger than associations with MI+revascularization (data not presented).

**All-cause mortality**—Measurable cTnT had a graded association with death (n=1,210; mean follow-up 9.9 years; Figure 2B) that persisted even in the fully adjusted model (HR 1.37, 95% CI 1.14–1.65 in Group 2; HR 3.96, 95% CI 3.21–4.88 in Group 5 [Table 2]). Associations between cTnT and mortality remained significant when men and women were evaluated separately (Supplemental Tables 2 and 3).

**HF Hospitalization**—Over a mean follow-up of 9.7 years, 668 HF hospitalizations occurred; ~3% (n=105) of individuals with undetectable cTnT had HF events compared with 25% (n=159) of individuals in the highest cTnT group. Detectable cTnT was associated with HF hospitalization even in the fully adjusted model; for the lowest and highest detectable cTnT categories, HRs were 1.48 (95% CI 1.14–1.92) and 5.95 (95% CI 4.47–7.92), respectively. Associations between cTnT and HF hospitalization remained significant at all detectable cTnT levels for men, while in women the association was significant at levels  $\geq 0.006$   $\mu\text{g/L}$  (Supplemental Tables 2 and 3).

Figure 2 shows cumulative incidence curves adjusted for age, gender, and race, and Figure 3 shows the continuous hazards functions, adjusted for ACRS components (model 2), of cTnT with total CHD events, all-cause mortality, and HF hospitalization. Overall, associations between cTnT and events were strongest for HF hospitalization, intermediate for all-cause mortality, and least robust for total CHD.

## Troponin T and Risk Prediction

Adding cTnT to ACRS reclassified 17.9% (n=1737) of the overall population for CHD risk prediction: 10.8% to lower risk and 7.1% to higher risk (Supplemental Table 4). For all CHD and hard CHD, adding cTnT (as a continuous variable) to an ACRS-based model significantly improved the AUC adjusted for optimism, NRI, clinical NRI, and IDI in the overall population and in women; in men, all parameters except for NRI for all CHD were significantly improved (Table 3). Improvements in statistical parameters for hard CHD prediction were generally better than for all CHD. Calibration of the models assessed by the Grønnesby–Borgan test indicated improved model fit with the addition of cTnT for hard CHD in the overall group and for all CHD in women. For mortality and HF hospitalization, adding cTnT to the base model (ACRS+BMI+LVH+creatinine) improved the AUC, NRI, clinical NRI, and IDI in all groups, and in general the improvements were greater in magnitude than for all CHD (no statistical comparisons performed). The receiver operator characteristics curves for the various outcomes are provided in the supplement (Supplemental Figure 1). In limited exploratory analyses, addition of either cTnT or NT-

proBNP appeared to be superior to adding hs-CRP for improving risk prediction for all outcomes (Supplemental Tables 5–7).

## DISCUSSION

In our analysis, cTnT measured with a new high-sensitivity assay had strong associations with CHD, mortality, and HF in a general population of middle-aged to older adults without CVD. The new assay detected cTnT in 66.5% of these individuals, the vast majority of whom had levels that would have been undetectable by the currently available, clinically used 4th-generation assay. Only 65 individuals (<1% of the ARIC population) had cTnT levels  $\geq 0.03$   $\mu\text{g/L}$  (i.e., detectable using the currently used assay), consistent with reported detectable levels in <1% of the general population<sup>5</sup> and 4.1% of the elderly population.<sup>6</sup>

Prior studies using this hs-cTnT assay have reported associations with adverse cardiovascular outcomes in individuals with chronic HF<sup>9</sup> and stable CHD,<sup>10</sup> and we now extend the findings to a general population aged 54–74 years. Among the outcomes tested, associations were strong with fatal CHD, all-cause mortality, and HF hospitalization. Of the CHD outcomes, the association with fatal CHD was strongest, and associations with nonfatal CHD events, including MI and revascularization, were weaker. However, relatively few fatal CHD events occurred; hence, when all CHD events were considered together, the association was weaker than with all-cause mortality and HF. Metrics of risk prediction, including AUC, NRI, and IDI, were also more robust for mortality and HF than for CHD (Table 3). In exploratory analyses, improvements in risk prediction for CHD, mortality, and HF when cTnT was added to risk prediction models were similar to those provided by NT-proBNP and larger than those for hs-CRP. In summary, our findings suggest that cTnT may be an important marker in the prediction of hard CHD, mortality, and HF.

Our results are concordant with those of 2 other population studies, the Cardiovascular Health Study (CHS)<sup>16</sup> and the Dallas Heart Study (DHS),<sup>17</sup> which were published after submission of this report; all 3 studies used the same assay. The ARIC results are most similar to those of CHS (average age ~10 years older than ARIC), in which 66% of participants had detectable cTnT with the new assay.<sup>16</sup> Although overall only 25% of DHS participants had detectable levels, >50% were aged <50 years; 56% of participants aged 60–65 had detectable levels.<sup>17</sup> All 3 studies found strong associations with mortality. The younger age and greater number of ARIC participants may have contributed to the larger NRI for mortality and HF than in CHS. The prevalence of detectable cTnT with the high-sensitivity assay is clearly distinct from the “sensitive” troponin I assay used by Blankenberg et al., which detected levels in <2% of population cohorts.<sup>8</sup>

The weaker association observed with nonfatal CHD events seems counter to findings in patients with acute coronary syndromes, in whom elevated cTnT level is a strong marker for risk for future MI.<sup>18,19</sup> However, small elevations in cTnT had no association with short-term risk for MI in asymptomatic individuals with stable CHD<sup>10</sup>; this, taken together with our findings, suggests that the risk from low detectable cTnT levels in asymptomatic subjects may be mediated through mechanisms other than or in addition to atherothrombosis. These mechanisms, which are not fully understood, may result from troponin release from cardiac myocytes due to asymptomatic ischemia, coronary microvascular dysfunction, apoptosis, or subclinical cardiac structural or functional abnormalities. Troponin I levels measured with a highly sensitive assay increased with reversible ischemia detected by nuclear perfusion imaging,<sup>20</sup> supporting the notion that subclinical ischemia could induce troponin release. However, in another study that used the hs-cTnT assay, cTnT levels did not change with exercise- or pharmacological stress-induced ischemia,<sup>21</sup> again suggesting that factors other than ischemia contribute



significantly to the risk associated with cTnT. Coronary microvascular dysfunction, which occurs in hypertension, diabetes, and LVH, is another potential cause for elevated circulating troponin levels<sup>22</sup> and may mediate, in part, the associations observed with HF events. Troponin is eliminated from the circulation via renal clearance; therefore, as glomerular filtration declines, troponin levels may increase. While this was true in ARIC, the associations with outcomes remained strong even after adjustment for eGFR. Therefore, the mechanisms underlying elevated troponin levels may be several, and it is unclear which mechanisms are related to the outcomes observed in our study.

## Clinical and Therapeutic Implications

The strength of the associations between cTnT and the various outcomes has several potential implications. First, troponin levels above 99<sup>th</sup> percentile values have been recommended for diagnosis of MI.<sup>1</sup> The 99<sup>th</sup> percentile value for cTnT in our study (0.03 µg/L) was far higher than that published by the manufacturer (0.014 µg/L), and 7% of the ARIC population had levels ≥0.014 µg/L. Januzzi et al.<sup>23</sup> reported (using the manufacturer-determined 99<sup>th</sup> percentile cutpoint) that the high-sensitivity assay had a lower positive predictive value (38%) than the conventional assay (67–72%), suggesting that false positives were higher with the high-sensitivity assay. If the 99<sup>th</sup> percentile identified in a population study such as ARIC were used, the estimated predictive value may be different. The diagnostic criteria may need to be revised to include both absolute cTnT level and rise and fall in cTnT level when the high-sensitivity assay is used.<sup>24</sup> Conversely, undetectable cTnT with the new assay may have superior negative predictive value, which will need further study.

cTnT modestly improves CHD risk prediction, and therapies such as statin and/or aspirin could hypothetically be considered for individuals reclassified as higher risk with cTnT. However, although statins reduce CHD events in patients with traditional risk factors such as hypertension,<sup>25</sup> statins have not shown benefit in some high-risk populations such as patients with HF<sup>26</sup> or end-stage renal disease.<sup>27</sup> Furthermore, a recent study in individuals with diabetes suggested that cTnT levels did not change significantly with either intensive or standard risk factor management.<sup>28</sup>

The stronger association of cTnT with death and HF than with CHD suggests that this biomarker predicts structural heart disease events to a greater extent than atherosclerotic events. Therefore, strategies aimed to prevent CHD may have smaller effects than strategies aimed to prevent progression of structural heart disease. As HF prevalence and incidence continue to increase, markers such as cTnT may conceivably identify high-risk individuals and allow early initiation of preventive strategies. Better understanding of the adverse mechanisms/process(es) that cTnT marks or mediates may be useful in targeting therapies, but ultimately clinical trials will be needed to examine if risk can be modified.

**Limitations**—Validated mortality and HF risk prediction models have not yet been developed in ARIC; therefore, we added BMI, creatinine, and LVH to the ACRS for use as the base model. However, AUCs were greater for both mortality (0.719) and HF (0.749) than for all CHD (0.715). We do not know if cTnT would have added to risk prediction with the addition of other biomarkers or clinical variables including echocardiography (which was not available) to the ACRS.

## Conclusion

cTnT measured with a novel highly sensitive cTnT assay was detectable in the majority of middle-aged individuals without prevalent CVD. Even slight elevations were strongly associated with death, especially CHD death, and HF hospitalization. Although there was an

association between cTnT and MI, it was less pronounced than that for the other endpoints assessed. Whether the risk for CHD, mortality, and HF hospitalization associated with measurable cTnT in individuals without prior CVD is modifiable is unknown and requires further study.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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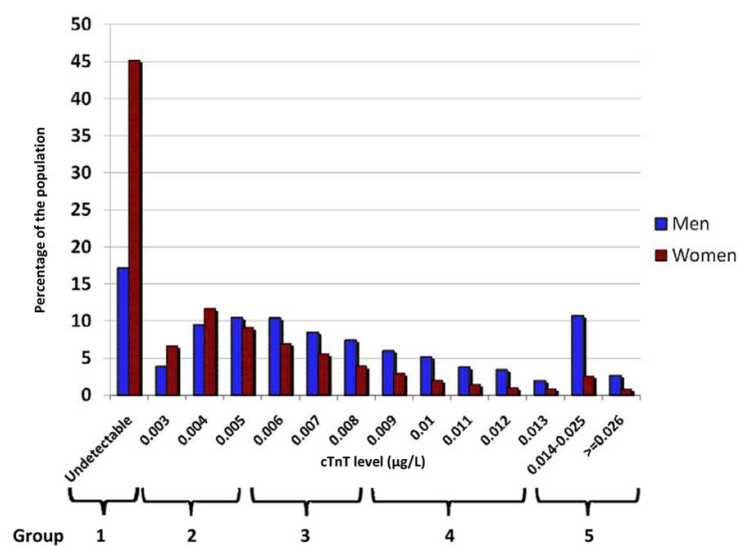
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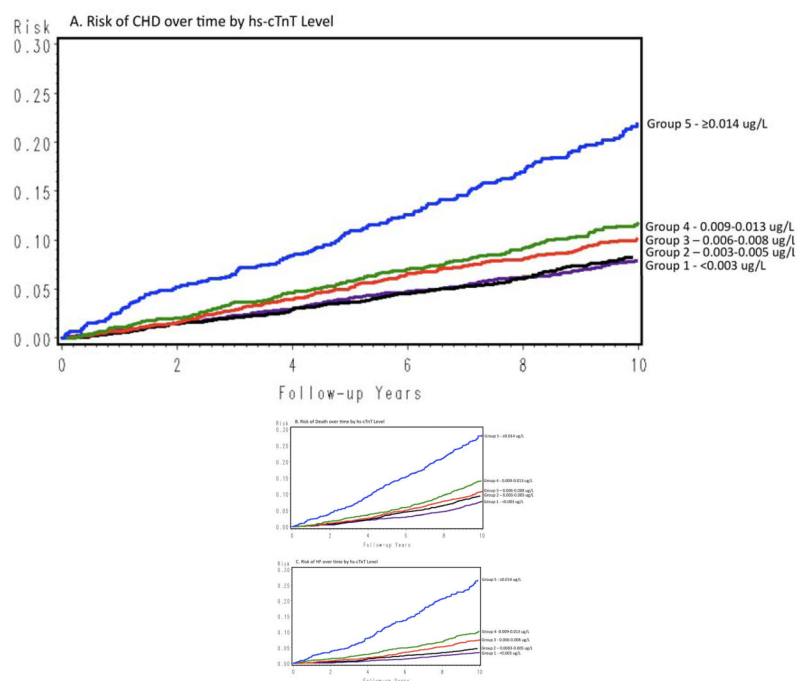
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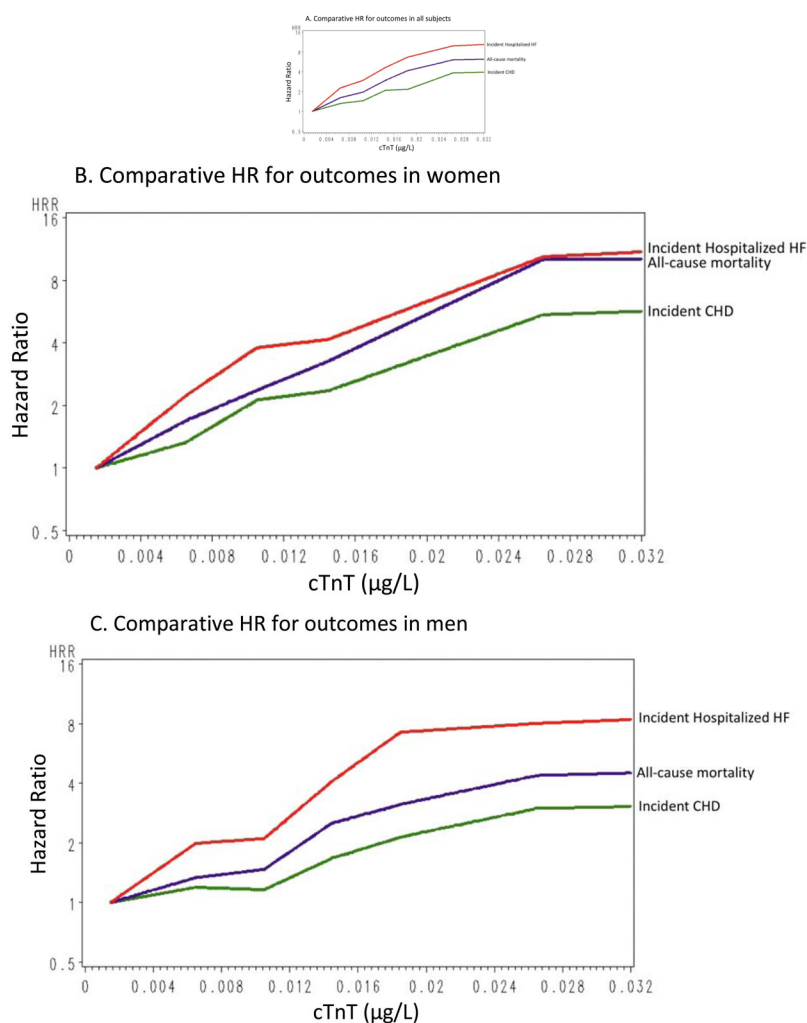
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**Figure 1.** Histogram showing gender-specific distribution of cardiac troponin T (cTnT) assay in the Atherosclerosis Risk in Communities (ARIC) sample.



**Figure 2.** Age-, race-, and gender-adjusted survival curves assessing the time to event across cTnT categories.



**Figure 3.** Continuous hazard functions of cTnT with total coronary heart disease, all-cause mortality, and heart failure hospitalizations for the overall group (A), in women (B), and in men (C), adjusted for age, race, gender (in the overall analyses), and traditional risk factors (total cholesterol, high-density lipoprotein cholesterol [HDL-C], systolic blood pressure, use of antihypertensive medications, smoking status, and presence of diabetes or the use of antidiabetic medications).

Table 1

Adjusted baseline characteristics\*

	Troponin T Group					P
	Group 1 (<0.003 µg/L) (n=3258)	Group 2 (0.003 µg/L to ≤0.005 µg/L) (n=2500)	Group 3 (0.006 µg/L to ≤0.008 µg/L) (n=1971)	Group 4 (0.009 µg/L to ≤0.013 µg/L) (n=1254)	Group 5 ≥0.014 µg/L (n=715)	
Demographic						
• Age, yrs	60.7	62.4	63.6	64.9	65.2	<0.0001
• Race, % white	78.2	80.7	78.2	76.6	70.4	<0.0001
• Gender, % male	20.9	37.7	52.8	63.6	73.7	<0.0001
• BMI, kg/m <sup>2</sup>	27.8	28.6	29.4	30.0	30.4	<0.0001
Medical History						
• Hypertension, %	38.7	42.	48.2	53.6	61.9	<0.0001
• Current smoking, %	22.0	13.0	10.0	10.0	11.2	<0.0001
• Former smoking, %	40.5	42.2	42.6	43.8	43.7	0.297
• Diabetes mellitus, %	9.8	12.4	17.3	21.1	35.4	<0.0001
• Systolic blood pressure, mm Hg	125.2	126.6	128.1	130.5	130.8	<0.0001
• Diastolic blood pressure, mm Hg	70.7	71.1	71.5	71.9	70.9	0.0088
Laboratory Data						
• Total cholesterol, mg/dL	203.6	202.1	201.1	200.6	197.3	0.0027
• HDL-C, mg/dL	51.8	51.0	50.2	49.6	48.7	<0.0001
• Triglycerides, mg/dL	137.8	139.1	142.8	149.5	157.1	0.0002
• eGFR, mL/min <sup>1.73</sup>	86.6	85.6	84.0	82.5	76.8	<0.0001
• Mean hs-CRP, mg/L <sup>‡</sup>	4.2	4.2	4.3	4.5	6.9	<0.0001
• Mean NT-proBNP, pg/mL <sup>§</sup>	70.1	92.4	112.4	149.9	530.7	<0.0001
Medications						
• Aspirin, %	53.9	53.8	53.3	54.9	57.9	0.310
• Antihypertensives, %	27.8	30.8	37.0	41.3	50.6	<0.0001



		Troponin T Group					P
		Group 1 (<0.003 µg/L) (n=3258)	Group 2 (0.003 µg/L to ≤0.005 µg/L) (n=2500)	Group 3 (0.006 µg/L to ≤0.008 µg/L) (n=1971)	Group 4 (0.009 µg/L to ≤0.013 µg/L) (n=1254)	Group 5 ≥0.014 µg/L (n=715)	
• Statins, %		8.8	8.2	7.9	9.9	10.4	0.143
LVH, %//		1.8	2.0	3.3	4.2	8.2	<0.0001

\* Adjusted for age, gender, and race

<sup>†</sup> eGFR calculated with the Chronic Kidney Disease Epidemiology Collaboration formula

<sup>‡</sup> Unadjusted median hs-CRP (mg/L): Group 1, 2.68; Group 2, 2.30; Group 3, 2.14; Group 4, 2.10; Group 5, 2.74; p<0.0001

<sup>§</sup> Unadjusted median NT-proBNP (pg/mL): Group 1, 59.00; Group 2, 61.65; Group 3, 61.60; Group 4, 71.30; Group 5, 98.00; p<0.0001

// LVH calculated by electrocardiographic Cornell Criteria

**Table 2**  
Cox proportional-hazards models for associations between troponin T and outcomes for overall cohort

	HR (95% CI)				
	Group 1 (<0.003 µg/L)	Group 2 (0.003 µg/L to ≤0.005 µg/L)	Group 3 (0.006 µg/L to ≤0.008 µg/L)	Group 4 (0.009 µg/L to ≤0.013 µg/L)	Group 5 ≥0.014 µg/L
ALL CHD					
	(n=3258, events=214)	(n=2500, events=205)	(n=1971, events=221)	(n=1254, events=171)	(n=715, events=172)
Model 1	Reference	1.06 (0.88, 1.29)	1.33 (1.09, 1.62)	1.50 (1.21, 1.86)	2.97 (2.38, 3.71)
Model 2	Reference	1.08 (0.89, 1.31)	1.31 (1.07, 1.59)	1.37 (1.10, 1.71)	2.46 (1.96, 3.08)
Model 3	Reference	1.07 (0.88, 1.30)	1.29 (1.06, 1.58)	1.34 (1.07, 1.67)	2.29 (1.81, 2.89)
HARD CHD (FATAL CHD + MI)					
	(n=3258, events=118)	(n=2500, events=104)	(n=1971, events=107)	(n=1254, events=81)	(n=715, events=117)
Model 1	Reference	1.02 (0.78, 1.33)	1.22 (0.93, 1.60)	1.34 (0.99, 1.82)	3.74 (2.81, 4.99)
Model 2	Reference	1.06 (0.81, 1.39)	1.26 (0.96, 1.67)	1.31 (0.96, 1.78)	3.28 (2.44, 4.42)
Model 3	Reference	1.05 (0.80, 1.38)	1.23 (0.93, 1.62)	1.23 (0.90, 1.68)	2.84 (2.09, 3.86)
ALL-CAUSE MORTALITY					
	(n=3258, events=217)	(n=2500, events=246)	(n=1971, events=248)	(n=1254, events=234)	(n=715, events=265)
Model 1	Reference	1.27 (1.05, 1.52)	1.45 (1.20, 1.76)	1.94 (1.59, 2.37)	4.34 (3.55, 5.29)
Model 2	Reference	1.39 (1.15, 1.67)	1.64 (1.35, 1.98)	2.13 (1.74, 2.60)	4.43 (3.61, 5.44)
Model 3	Reference	1.37 (1.14, 1.65)	1.60 (1.32, 1.94)	2.05 (1.68, 2.51)	3.96 (3.21, 4.88)
HF HOSPITALIZATION					
	(n=3158, events=105)	(n=2413, events=124)	(n=1877, events=147)	(n=1188, events=130)	(n=640, events=159)
Model 1	Reference	1.45 (1.12, 1.88)	2.21 (1.71, 2.86)	3.07 (2.34, 4.04)	8.61 (6.57, 11.28)
Model 2	Reference	1.51 (1.16, 1.96)	2.24 (1.73, 2.90)	2.84 (2.16, 3.74)	7.00 (5.29, 9.25)
Model 3	Reference	1.48 (1.14, 1.92)	2.17 (1.67, 2.81)	2.68 (2.03, 3.53)	5.95 (4.47, 7.92)

Model 1: adjusted for age, gender, and race

Model 2: adjusted for ARIC CHD Risk Score (Model 1 + total cholesterol, HDL-C, systolic blood pressure, antihypertensive medication use, smoking status, and presence of diabetes defined as fasting blood glucose >126 mg/dL or antidiabetic medication use)

Model 3: Model 2 + eGFR + hs-CRP + NT-proBNP

Table 3

Metrics for improvement in risk prediction with addition of cardiac troponin T (cTnT)

	Overall		Women		Men	
	Base model	Base model + cTnT	Base model	Base model + cTnT	Base model	Base model + cTnT
<b>HARD CHD (FATAL CHD + MI)*</b>						
Adjusted AUC (95% CI)	0.710 (0.691, 0.732)	0.724 (0.708, 0.747)	0.692 (0.667, 0.725)	0.718 (0.695, 0.753)	0.688 (0.663, 0.720)	0.701 (0.679, 0.735)
Difference in AUC (95% CI)		0.014 (0.008, 0.024)		0.027 (0.014, 0.046)		0.013 (0.004, 0.028)
G-B statistic	7.02 (p=0.64)	6.78 (p=0.66)	12.55 (p=0.18)	12.72 (p=0.18)	8.51 (p=0.48)	12.83 (p=0.17)
IDI (95% CI)		0.032 (0.021, 0.048)		0.043 (0.026, 0.071)		0.028 (0.015, 0.052)
NRI (95% CI)		10.1% (3.8%, 15.9%)		19.2% (6.7%, 25.2%)		7.2% (1.6%, 19.1%)
NRI in intermediate-risk group (95% CI)		22.4% (13.7%, 33.4%)		42.0% (20.5%, 57.1%)		15.2% (6.9%, 29.0%)
<b>ALL CHD*</b>						
Adjusted AUC (95% CI)	0.715 (0.702, 0.733)	0.724 (0.711, 0.742)	0.691 (0.670, 0.719)	0.704 (0.684, 0.730)	0.660 (0.643, 0.684)	0.673 (0.656, 0.699)
Difference in AUC (95% CI)		0.009 (0.005, 0.015)		0.012 (0.004, 0.024)		0.013 (0.005, 0.024)
G-B statistic	14.24 (p=0.11)	19.87 (p=0.02)	17.88 (p=0.04)	6.49 (p=0.69)	9.74 (p=0.37)	14.79 (p=0.10)
IDI (95% CI)		0.021 (0.014, 0.032)		0.025 (0.014, 0.040)		0.021 (0.012, 0.037)
NRI (95% CI)		4.5% (0.4%, 8.8%)		6.9% (1.1%, 17.1%)		2.0% (-0.5%, 9.6%)
NRI in intermediate-risk group (95% CI)		11.7% (6.9%, 17.9%)		19.8% (9.9%, 32.1%)		6.9% (2.8%, 14.3%)
<b>TOTAL MORTALITY†</b>						
Adjusted AUC (95% CI)	0.719	0.740	0.692	0.715	0.712	0.733
Difference in AUC (95% CI)		0.021 (0.014, 0.028)		0.024 (0.014, 0.034)		0.021 (0.013, 0.031)
G-B statistic	11.8 (p=0.23)	8.8 (p=0.46)	12.6 (p=0.18)	8.0 (p=0.53)	15.9 (p=0.07)	4.1 (p=0.91)
IDI (95% CI)		0.037 (0.027, 0.047)		0.035 (0.022, 0.050)		0.037 (0.025, 0.052)
NRI (95% CI)		10.7% (4.3%, 13.1%)		10.0% (4.2%, 16.9%)		6.1% (1.4%, 13.0%)
NRI in intermediate-risk group (95% CI)		21.4% (14.4%, 25.9%)		21.0% (13.0%, 29.4%)		18.5% (9.9%, 26.3%)
<b>HF HOSPITALIZATION‡</b>						
Adjusted AUC (95% CI)	0.749	0.777	0.741	0.769	0.736	0.772
Difference in AUC (95% CI)		0.028 (0.019, 0.037)		0.028 (0.019, 0.042)		0.036 (0.022, 0.053)
G-B statistic	11.5 (p=0.24)	10.4 (p=0.32)	10.1 (p=0.34)	9.1 (p=0.42)	9.2 (p=0.42)	2.3 (p=0.99)

	Overall		Women		Men	
	Base model	Base model + cTnT	Base model	Base model + cTnT	Base model	Base model + cTnT
IDI (95% CI)		0.044 (0.032, 0.058)		0.038 (0.023, 0.055)		0.057 (0.036, 0.085)
NRI (95% CI)		15.4% (8.4%, 20.6%)		16.7% (6.4%, 22.4%)		15.9% (8.0%, 27%)
NRI in intermediate-risk group (95% CI)		30.9% (22.1%, 40.0)%		31.2% (18.0%, 43.6%)		30.7% (20.1%, 49.5%)

AUC = area under the receiver operating characteristic curve; G-B statistic = Grønnesby – Borgan goodness-of-fit statistic; IDI = integrated discrimination improvement; NRI = net reclassification improvement

\* Base model for CHD: model based on ARIC Coronary Risk Score (age, gender, race, total cholesterol, HDL-C, systolic blood pressure, use of antihypertensive medications, smoking status, and presence of diabetes defined as fasting blood glucose >126 mg/dL or use of antidiabetic medications).

† Base model for total mortality and HF: ARIC Coronary Risk Score + BMI + presence of LVH on ECG + serum creatinine.